

**Table S1: General characteristics of GGI networks**

<b>Organism</b>	<b>Network type</b>	<b>#Nodes</b>	<b>#Interactions</b>	<b>&lt;K&gt;</b>	<b>γ</b>	<b>C</b>	<b>C<sub>rand</sub></b>	<b>l</b>	<b>l<sub>rand</sub></b>
Human	<b>PPI network</b> (HIPPIE, <sup>1</sup> Cutoff: 0.68)	13987	130761*2	18.7	1.50	0.051	0.0013	3.34	3.60
	<b>PPI network</b> (HIPPIE, <sup>1</sup> Cutoff: 0.77)	9656	45842*2	9.5	1.67	0.071	0.0012	3.97	4.33
	<b>PPI network</b> (HPRD)	8621	32342*2	7.5	1.81	0.055	0.0008	4.26	4.7
	<b>GC network</b> ( <sup>2</sup> Cutoff: 0.4)	13123	794265*2	121	1.19	0.457	0.0092	3.23	2.32
Mouse	<b>PPI network</b> ( <sup>1</sup> Cutoff: 0.68)	4633	9921*2	4.3	1.49	0.029	0.0010	4.36	6.04

**Abbreviations:**

PPI: Protein-Protein Interaction; GC: Gene Co-expression;

<sup>1</sup>Confidence score; <sup>2</sup>Pearson correlation coefficient;

#Nodes: Number of nodes; #Interactions: Number of interacting links; &lt;K&gt;: Average connectivity (degree) of network;

γ: Degree exponent of scale-free network; C: Clustering coefficient of network; l: Average path length of network;

C<sub>rand</sub>, l<sub>rand</sub>: Clustering coefficient and average path length of random network with the same numbers of nodes and interactions.